



SEQ ID NO: 92 DR01006
 SEQ ID NO: 93 DR27006
 SEQ ID NO: 94 DR03006
 SEQ ID NO: 95 DR11006
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 SEQ ID NO: 97 DR16006
 SEQ ID NO: 98 DR19006
 SEQ ID NO: 99 DR07006
 SEQ ID NO: 100 DR16006
 SEQ ID NO: 101 DR20006
 SEQ ID NO: 102 DR25006
 SEQ ID NO: 103 DR20006
 SEQ ID NO: 104 DR21006
 SEQ ID NO: 105 DR09006
 SEQ ID NO: 106 DR17006
 SEQ ID NO: 107 DR13006
 SEQ ID NO: 108 DR02006

C-----TCGAG---TCTGGGGGAGG
 C-----TCGAG---TCTGGGGGAGG
 C-----AGGTGA----AACTGCTCGAG---TCTGGAGGAGG
 C-----TCGAG---TCTGGGGGAGG
 C-----AGGTGA----AACTGCTCGAG---TCTGGGGGAGG
 C-----TCGAG---TCTGGAGGAGG
 C-----TCGAG---TCTGGGGGAGG
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 C-----TCGAG---TCTGGGGGAGG
 C-----TCGAG---TCTGGGGGAGG
 C-----TCGAG---TCTGGGGGAGG
 C-----TCGAG---TCAAGGGGGAGG
 C-----TCGAG---TCTGGGGGAGG
 C-----TCGAG---TCTGGGGGAGG
 C-----AGGTGA----AACTGCTCGAG---TCTGGGGGAGG
 C-----TCGAG---TCTGGGGGAGG
 C-----TCGAG---TCAAGGGGGAGG
 C-----TCGAG---TCTGGGGGAGG
 C-----TCGAGTCAGGTGTCCGGTCTGATGTGCAGCTGGTGGCGTCTGGGGGAGG

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ATCGGTGCAGGCTGGAGGGTCTCTGAGACTCTC--GTGCG-CAGCCTCTG
 CTCGGTGCAGGCTGGAGGGTCTCTGAGACTCTCCTGTGCATTTCTCTA
 CTCGGTGCAGGACTGGAGGGATCTCTGAGACTCTCCTGTGCAGT--C-TCTG
 GTCGGTGCAGGCTGGAGGGTCTCTGAGACTCTCCTGTAAATGT--C-TCTG
 GTCGGTGCAGGCTGGAGGGTCTCTGAGACTCTCCTGTAAATGT--C-TCTG
 CTCGGCGCAGGCTGGAGGGATCTCTGAGACTCTCCTGTGCAGC--CCACGG
 CTCGGTTAGGGCTGGAGGGTCCCTTAGACTCTCCTGTGCAGC--C-TCTG
 CTCGGTGCAGGGTGGAGGGTCTCTGAGACTCTCCTGTGCAA--TCTCTG
 CTCGGTGCAGGCTGGAGGGTCTCTGAGACTCTCCTGTACAG--GCTCTG
 CTCGGTACAGGTTGGAGGGTCTCTGAGACTCTCCTGTGTAG--CCTCTA
 CTCGGTACAAACTGGAGGGTCTCTGAGACTCTTGC--AAATCTCTG
 CTCGGTGCAGGCTGGAGGGTCTCTGAGACTCTCCTGTG--TAGCCTCTG
 CTCGGTGCAGGTTGGAGGGTCTCTGAAACCTCTCCTGTAAAAT--CTCTG
 CTCGGTGCAGGCTGGGGGGTCTCTGACACTCTTGTG--TATAACAC--
 CTCGGTCCAACCTGGAGGGATCTCTGACACTCTCCTGTACAGTT--TCTG
 CTCGGTGGAGGGCTGGAGGGTCTCTGAGACTCTCCTGTACAG--CCTCTG
 CTCGGTGCAGGCTGGAGGCTCTCTGAGACTCTCCTGTACAG--CCTCTG

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GA--TACAGTAATT---GTCCCCTCACTTG-GAGCTGGTATGCCAGTT
 AA--TATATGCCCTT---GCACCTACGACAT-GACCTGGTACCGCCAGGCT
 GA--TTCTCCTTTA---GTACCAAGTTGTAT-GGCCTGGTTCCGCCAGGCT
 GC--TCTCCCAGTA---GTACTTATTGCCT-GGGCTGGTTCCGCCAGGCT
 GC--TCTCCCAGTA---GTACTTATTGCCT-GGGCTGGTTCCGCCAGGCT
 GA--TTCCGC-TCA---ATGGTTACTACAT-CGCCTGGTTCCGTCAGGCT
 AC--TACACCATCA---CTGATTATTGCAT-GGCCTGGTTCCGCCAGGCT
 GA--TACACGTACG---GTAGCTTCTGTAT-GGGCTGGTTCCGCCAGGCT
 GA--TTCCCCTATA---GTACCTTCTGTCT-GGGGTGGTTCCGCCAGGCT
 CT--CACACCGACA---GTAGCACCTGTAT-AGGCTGGTTCCGCCAGGCT
 GA--TTGACTTTG---ATGATTCTGACGT-GGGGTGGTACCGCCAGGCT
 GA--TTCAATTTCG---AAACTTCTCGTAT-GGCCTGGTACCGCCAGACT
 GAGGTACCCAGATCGTGTCTAAATCTTGGCCTGGTTCCGCCAGGCT
 -----CAACGATACTGGGACCA-----TGGGATGGTTCCGCCAGGCT
 --GGGCCACCACA---GTGACTACAGTATTGGA-TGGATCCGCCAGGCT
 G-----ATACGTAT-CCT---CTATGGCCTGGTTCCGCCAGGTT
 GAGA---CAGTTCACTAGATT--TGCCATGTCTGGTTCCGCCAGGCT

FIG. 7A